## SCORE Search Results Details for Application 10537864 and Search Result 20080418 161306 us-10-537-864a-2.p2n.rni.

Score Home Page

Retrieve Application

SCORE System Overview

SCORE

Comments / Suggestions

Appendix A

This page gives you Search Results detail for the Application 10537864 and Search Result 20080418\_161306\_us-10-537-864a-2.p2n.mi.

Go Back to previous page

GenCore version 6.2.1 Copyright (c) 1993 - 2008 Biocceleration Ltd.

OF Drotein - nucleic search, using frame plus p2n model

April 18, 2008, 19:29:58; Search time 1023 Seconds (without alignments) 1055.383 Million cell updates/sec

Tinle: U8-10-537-864A-2
Perkent score: 1442
Segmence: 1 AARENVAVAPUSLALSQVVC......LLDSPQQATYPSAFKVLDID 275

Mgapop 10.0 . Xgapext 0.5 Ygapop 10.0 . Ygapext 0.5 Fgapop 6.0 . Fgapext 7.0 Delop 6.5 . Delext 7.0

5376\$16 segs, 1963011933 residues

Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 2000000000

Post processing: Minimum Match 90% Maximum Match 100% Listing first 45 summaries

Command line parameters:
-TAN-4 -NODEL-frame, p2n. model -DRV-SOFT
-CA-Abs/JASSNEES spool/US105:7864/rumat\_18042008\_16:306\_17814/app\_query\_fasta\_1
-DR-SSENEED PALEMIA\_NODEL-FRAMES\_NODEL-FRAMES\_0.0005
-TRESSENEED PALEMIA\_NODEL-FRAMES\_NODEL-FRAMES\_0.0005
-TRESSENEED PALEMIA\_NODEL-FRAMES\_NUMBER\_0.0005
-TRESSENEED PALEMIA\_NODEL-FRAMES\_NUMBER\_0.0005
-TRESSENEED NUMBER\_0.0005
-TRESSENE

Issued\_Patents\_Na.\*

1: /ABSS/Data/CRF/ptodata/Z/ina/1\_COMB.seq.\*

2: /ABSS/Data/CRF/ptodata/Z/ina/5\_COMB.seq.\*

3: /ABSS/Data/CRF/ptodata/Z/ina/6a\_COMB.seq.\*

3: /ABSS/Data/CRF/ptodata/Z/ina/6a\_COMB.seq.\*

5: /ABSS/Data/CRF/ptodata/Z/ina/7a\_COMB.seq.\*

5: /ABSS/Data/CRF/ptodata/Z/ina/7a\_COMB.seq.\*

7: /ABSS/Data/CRF/ptodata/Z/ina/7a\_COMB.seq.\*

7: /ABSS/Data/CRF/ptodata/Z/ina/f\_COMB.seq.\*

9: /ABSS/Data/CRF/ptodata/Z/ina/FCTUS\_COMB.seq.\*

9: /ABSS/Data/CRF/ptodata/Z/ina/FCTUS\_COMB.seq.\*

10: /ABSS/Data/CRF/ptodata/Z/ina/FE\_COMB.seq.\*

11: /ABSS/Data/CRF/ptodata/Z/ina/Backfiles1.seq.\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

Description Query Score Match Length DB ID 1317 91.3 954 3 US-06-996-139-12 Sequence 12, Appl

```
Enquence 12, Appl
Sequence 2, Appl
                                                                                                                                                                                                                                    US-08-995-659-12
US-09-215-649A-12
US-09-877-740-12
US-09-877-800-12
US-09-466-496-13
                                                                                                                                                      994 3
964 3
954 3
954 3
954 3
954 3
954 3
954 3
954 3
954 5
984 3 03-09-466-494-12
984 3 03-09-466-494-13
984 3 03-09-471-281-12
984 3 03-09-471-281-12
984 3 03-09-487-480-12
984 3 03-09-487-480-12
984 3 03-09-887-944-5
984 5 03-09-887-944-5
984 6 03-10-465-878-12
2271 3 03-09-481-2812-3
2271 3 03-09-481-381-38
2271 3 03-09-981-381-38
                                                                     91.3
91.3
91.3
91.3
91.3
91.3
91.3
91.3
```

## Appendix A

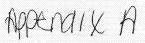
## ALIGNMENTS

```
Alignment Scores:
Pred. No.:
Score:
Percent Similarity:
Mest Local Similarity:
Obery Match:
DB:
                                                                     Length:
Matches:
Conservative:
Mismatches:
                                           91.3%
                                                                     Indels:
Gaps:
  08 \cdot 10 \cdot 537 \cdot 864 A \cdot 3 \cdot (1 \cdot 278) \times 98 \cdot 98 \cdot 996 \cdot 138 \cdot 12 \cdot (1 \cdot 984)
```

```
06
                                              667 CDACATCATGAAACTTCAGGAGACCTAGCTACAGAGTATCTTCAACTAATGGTGTACGTC 726
  Qv
                                            201 TorbysThrSerfleLyalleProSerSerHisThrLeuMetLyaClyClySerThrLys 220
                                             727 ACTARARCENSCRICARARTECCARGITYTTCATRCCCTGATGARAGGARGGACGACCARG 76K
  ďĊ
                                           221 TyrtrpSerGlyAsnSerGluRheHisPheTyrSer]leAsmValGlyGlyRhePheLyw 245
  Qy
  Dis
                               Dès
                                           261 GlnAspAlathityiyhesiyaiaphetysVelLeuAspIleAsp 275
  Ov
  Db
 RESULT 1:
US-09-588-459-12
; Sequence 12. Application US/09688459
; Patent No. 5838262
; GENERAL INFORMATION:
         OUR -58-459-12
Acquence 12. Application US/OPERESS
Areat No. 538252
GORREAL INFORMATION.
APPLICANT: Galibert Laurent
APPLICANT: Macaskovsky Engene
TITLE OF INVERTION Liquand for Receptor Activator of NF-kappaB
RUMBER OF SEQUENCES: 19
CORRESORDERCE ADDRESS:
ADORRSORS: Ismunez Corporation, Law Department
STREET: SI University Street
CITY: Seattle
STATE: MA
COUNTRY: USA
ZIP: 98101
COMPUTER READBLE FORM:
MEDIUM TYPE: PLoppy disk
COMPUTER READBLE FORM:

CHARLING STREET APPLICATION DATA:
APPLICATION RUMBER: US/OP/688,459
FILING DATE:
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION MUMBER: US/OP/688,459
FILING DATE: OT MARCH 1997
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/OP/688,459
FILING DATE: OT MARCH 1997
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/OP/72,330
FILING DATE: OT MARCH 1997
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/OP/72,330
FILING DATE: OT MARCH 1997
CLASSIFICATION:
MINERAL SCHOOL:
ATTORNEY/ACCENT INFORMATION:
NAME: PEXIST, PALYICLE AND
REFERENCE/COCKET MUMBER: 2852
TELEFANCE: COMPUTER COM
  NAME/KEY. CDS
1.OCATION: 1.951
US-09-E58-459-12
Alignment Scores:
Pred No.: 137e-156
Score: 137.00
Percent Similarity: 94.54
Scot Local Similarity: 90.59
Obery Match: 91.34
Db: 3
                                                                                                                                                                            Laugth:
Matches:
Conservative:
Mismatches:
Indels:
Gaps:
                                                                                                                                                                                                                                                 954
249
13
15
0
```

US-10-537-864A-2 (1-275) x US-09-688-459-12 (1-954)



```
i AlaalaSerArgSerValAlaValAlaFheixeuSlyLeuSlyLeuGlyCinValValCya 10
                                                                                                                                                                                                 Appendix A
                      127 GCCCCTCCCGCTCCATGTTCCTGGCCCTCCTCCGCGCTGGGCTGGGCTGGGCCAGGTTGTCTGC 186
  Sis
                      21 ServalalaLeuPhelevTyrPheargalaGlnMetaspProAsnArgIleSerGluAsp 40
  os
                      41 AspturniscysileäsnärgilePhelyslaudisciluäsnälaäspheuSinäspthr 60
   QÝ
  ಶಾಚ
                     ٥у
   oh:
   Qy
                       81 PheargalaalavalGinLyvGiuLeuGinHisileValargserGinHisilevangala 100
                      367 TTTCRAGGAGCTGTGCAAAAGGAATTACAACATATCGTTGGATCACAGCACATCAGAACA 426
   ಐಟ
                      Qy
   66
                      121 GlnBroPheAlaHisLeuThrIleAsnAlaThrAspIleProS#rGlySerHisLysVal 140
  Qy
                      487 CASECTITICCTCATCTCACTATTAATCCCACCGACCCCCATCTGGTTCCCCATAAGTG 548
   dø
                      141 SerteuSerSerTrpTyrHisaspArgGiyTrpAlaLysIleSerAsnMetThrFheSer 160
                      rah-
   Qy
                      161 AsnGlyLysLev(!eVslAsnGlnAspGlyPheTyrPheLeuTyrAlsAsn1leCysPhe 180
                      607 ANTOGRANACIRATINATICAGGATINOTTYATTACCTGTATTOCCRACATTINCTT 666
   Db
                      Ųу
   Dis
                      201 ThrtysThxSsr[]eliys1leProScrScrHisThrLeuMerLysG]yGlySerThrLys 220
   Ov
                      221 TyrTrpSerGlyAsnSerStoPhonisensTyrSerTheAsnValStyGlyPhePhelys 240
                      787 TATTNEFCNEGGAATTCTGAATTCCATTTTATTCCATAAACGTTGGTKGATTTTTTAAG 846
   ob
                      241 LeuArgSerOlyGluGluHeSerHeGiuVelSerAsnProSerLeuLeuAspProAsp 260
   Qy.
                      847 Tracquictggagasgaaatcascarcgaggictccaaccourtectractssarccogar 806
                     261 GlnaspalathrtyrPheGlyAlaPhelysVallenaspileasp 275
   Qy
NESULT 12

NESULT 12

(See 39 - 957 - 944 - 5)

(Sequence S. Application US/09957944

(Potent No. 688398)

(GENERAL INFORMATION:

APPLICANT: Dough 11, William C.

(TITLE OF INVENTION: SERENIMA ASSAYS FOR AGONISTS AND ANTAGONISTS OF RECEPTOR ACTIVATOR OF TITLE OF INVENTION: NF. ADPAIL AND ANTAGONISTS OF RECEPTOR ACTIVATOR OF TITLE OF INVENTION: NF. ADPAIL AND ANTAGONISTS OF RECEPTOR ACTIVATOR OF TITLE OF INVENTION: NF. ADPAIL AND ADPAIL AND ANTAGONISTS OF RECEPTOR ACTIVATOR OF TITLE OF INVENTION: NF. ADPAIL AND A
   Db
  Alignment Scores:
Fred So.:
Schree
Percent Similarity:
New Local Similarity:
Query Match.
ma.
                                                                                 Length:
Katches:
Conservative:
Mismatches:
Indels:
   US-10-537-864A-2 (1-275) x US-09-957-944-5 (1-954)
```

http://es/ScoreAccessWeb/GetItem.action?AppId=10537864&seqId=09323b678070341f&... 5/15/2008